



CGCTCAGGATATGACTTCGCTGCTAGAGGATCCGAGCCCGCTGCTATTATATAGTCGATCGAIC
TTCTCTATATCCCTGCTGCTAGAGGATCCGAGCCCGCTGCTATTATATAGTCGATCGAIC
CCGCTCAGGATATGACTTCGCTGCTAGAGGATCCGAGCCCGCTGCTATTATATAGTCGATCGAIC
CACTTACTAGCAGCTTCTGCTAGAGGATCCGAGCCCGCTGCTATTATATAGTCGATCGAIC

Entrez

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Nucleotide

Protein

Genome

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for ACP-CoA transacylase

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Entrez Nucleotide



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Links

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Haemophilus influenzae Rd KW20 section 16 of 163 of the complete
genome
gi|3212182|gb|U32701.1|[3212182]

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☐ 2: [U91631](#)

Links

Related resources
BLAST

Pseudomonas aeruginosa PlsX protein homolog (plsX) gene, partial cds; and
malonyl-CoA:acyl carrier protein transacylase (fabD), 3-oxoacyl-acyl
carrier protein reductase (fabG), acyl carrier protein (acpP), and 3-oxoacyl-
acyl carrier protein synthase II (fabF) genes, complete cds
gi|2738151|gb|U91631.1|PAU91631[2738151]

Reference sequence
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Vibrio harveyi malonyl-CoA:ACP transacylase (fabD) gene, partial cds, and
3-ketoacyl-ACP reductase (fabG), acyl carrier protein (acpP), 3-ketoacyl-
ACP synthase II (fabF) and aminodeoxychorismate lyase (pabC) genes,
complete cds
gi|1173839|gb|U39441.1|VHU39441[1173839]

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Links

Sorangium cellulosum acyl-CoA dehydrogenase (sorE), beta-hydroxybutyryl-CoA dehydrogenase (sorD), putative methoxymalonyl-CoA synthase (sorC), reductase (sorR), soraphen polyketide synthase A (sorA), soraphen polyketide synthase B (sorB), O-methyltransferase (sorM), beta-mannanase, and xylanase-arabinofuranosidase bifunctional enzyme genes, complete cds; and unknown genes
gi|13346872|gb|U24241.2|SCU24241[13346872]

☐ 2: [U37088](#)

Links

Simmondsia chinensis beta-ketoacyl-CoA synthase mRNA, complete cds
gi|1045613|gb|U37088.1|SCU37088[1045613]

☐ 3: [U75685](#)

Links

Mycobacterium bovis acyl-CoA synthase gene, complete cds
gi|1658530|gb|U75685.1|MBU75685[1658530]

☐ 4: [S73733](#)

Links

ACBP/DBI=acyl CoA-binding protein (ACBP)/diazepam-binding inhibitor (DBI)-endozepine homolog [Anas platyrhynchos=ducks, mallard, Genomic, 5191 nt]
gi|765222|bbm|351962|bbs|156494|gb|S73733.1|S73733[765222]

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Links

Garcinia mangostana acyl-ACP thioesterase (FatB1) mRNA, nuclear gene encoding chloroplast protein, complete cds
gi|1930080|gb|U92878.1|GMU92878[1930080]

☐ 2: U92877

Links

Garcinia mangostana acyl-ACP thioesterase (FatA2) mRNA, nuclear gene encoding chloroplast protein, complete cds
gi|1930078|gb|U92877.1|GMU92877[1930078]

☐ 3: U92876

Links

Garcinia mangostana acyl-ACP thioesterase (FatA1) mRNA, nuclear gene encoding chloroplast protein, complete cds
gi|1930076|gb|U92876.1|GMU92876[1930076]

☐ 4: U56104

Links

Cuphea wrightii acyl-ACP thioesterase (FatB2) mRNA, complete cds
gi|1336007|gb|U56104.1|CWU56104[1336007]

☐ 5: U56103

Links

Cuphea wrightii acyl-ACP thioesterase (FatB1) mRNA, complete cds
gi|1336005|gb|U56103.1|CWU56103[1336005]

☐ 6: U38189

Links

Cuphea palustris thioesterase (FatB2) mRNA, complete cds
gi|1215719|gb|U38189.1|CPU38189[1215719]

☐ 7: U38188

Links

Cuphea palustris thioesterase (FatB1) mRNA, complete cds
gi|1215717|gb|U38188.1|CPU38188[1215717]

☐ 8: U31813

Links

Cinnamomum camphora acyl-ACP thioesterase mRNA, complete cds
gi|1143155|gb|U31813.1|U31813[1143155]

☐ 9: U17076

Links

Cuphea hookeriana 16:0-ACP thioesterase preprotein (Ch FatB1) mRNA,

complete cds
gi|758700|gb|U17076.1|CHU17076[758700]

☐ 10: U17098

Links

Brassica rapa acyl-ACP thioesterase Br FatA1 (FatA1) mRNA, complete
cds

gi|595956|gb|U17098.1|BRU17098[595956]

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